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Date: 6/20/2006

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 **PALM INTRANET**

## Inventor Information for 09/349954

Inventor Name	City	State/Country
HAYWARD, NICHOLAS KIM	PADDINGTON	AUSTRALIA
WEBER, GUNTHER	STOCKHOLM	SWEDEN
GRIMMOND, SEAN	TARINGA	AUSTRALIA
NORDENSKJOLD, MAGNUS	STOCKHOLM	SWEDEN
LARSSON, CATHARINA	STOCKHOLM	SWEDEN

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## Inventor Information for 08/569063

Inventor Name	City	State/Country
ERIKSSON, ULF	BALSTA	SWEDEN
OLOFSSON, BIRGITTA	SUNDBYBERG	SWEDEN
ALITALO, KARI	HELSINKI	FINLAND
PAJUSOLA, KATRI	HELSINKI	FINLAND

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38	92.2	8.4	495	4	US-09-244-583-25	Sequence 25, Appl
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## ALIGNMENTS

## RESULT 1

US-09-949-016-1545

; Sequence 1545, Application US/09949016

; Patent No. 6812339

; GENERAL INFORMATION:

; APPLICANT: VENTER, J. Craig et al.

; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

; TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

; FILE REFERENCE: CL001307

; CURRENT APPLICATION NUMBER: US/09/949,016

; CURRENT FILING DATE: 2000-04-14

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## SUMMARIES

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## ALIGNMENTS

## RESULT 1

US-09-949-016-1545

; Sequence 1545, Application US/09949016

; Patent No. 6812339

; GENERAL INFORMATION:

; APPLICANT: VENTER, J. Craig et al.

; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

; TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

; FILE REFERENCE: CL001307

; CURRENT APPLICATION NUMBER: US/09/949,016

; CURRENT FILING DATE: 2000-04-14

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## ALIGNMENTS

## RESULT 1

US-08-469-427A-11

; Sequence 11, Application US/08469427A

; Patent No. 5607918

; GENERAL INFORMATION:

; APPLICANT: Eriksson, Ulf

; APPLICANT: Olofsson, Birgitta

; APPLICANT: Alitalo, Kari

; APPLICANT: Pajusola, Katri

; TITLE OF INVENTION: VASCULAR ENDOTHELIAL GROWTH FACTOR-B AND

; TITLE OF INVENTION: DNA CODING THEREFOR



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44	92.2	10.7	516	4	US-09-574-708A-3	Sequence 3, Appli
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## ALIGNMENTS

## RESULT 1

US-09-949-016-1545

; Sequence 1545, Application US/09949016

; Patent No. 6812339

; GENERAL INFORMATION:

; APPLICANT: VENTER, J. Craig et al.

; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

; TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

; FILE REFERENCE: CL001307

; CURRENT APPLICATION NUMBER: US/09/949,016

; CURRENT FILING DATE: 2000-04-14

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14	334	36.7	591	2	US-08-569-063C-6	Sequence 6, Appli
15	334	36.7	591	3	US-08-851-896-6	Sequence 6, Appli
16	329.8	36.2	7386	4	US-09-949-016-13287	Sequence 13287, A
17	312	34.3	565	1	US-08-469-427A-4	Sequence 4, Appli
18	312	34.3	565	2	US-08-609-443B-4	Sequence 4, Appli
19	312	34.3	565	2	US-08-569-063C-4	Sequence 4, Appli
20	312	34.3	565	3	US-08-851-896-4	Sequence 4, Appli
21	273.8	30.1	886	1	US-08-469-427A-1	Sequence 1, Appli
22	273.8	30.1	886	2	US-08-609-443B-1	Sequence 1, Appli
23	273.8	30.1	886	2	US-08-569-063C-1	Sequence 1, Appli
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25	239.4	26.3	405	1	US-08-469-427A-8	Sequence 8, Appli
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30	71.4	7.8	444	3	US-09-392-932-6	Sequence 6, Appli
31	71.4	7.8	444	4	US-09-574-708A-1	Sequence 1, Appli
32	71.4	7.8	444	4	US-09-392-931-1	Sequence 1, Appli
33	71.4	7.8	456	5	PCT-US95-10973A-88	Sequence 88, Appl
34	71.4	7.8	467	5	PCT-US95-10973A-86	Sequence 86, Appl
35	71.4	7.8	473	3	US-08-718-904-1	Sequence 1, Appli
36	71.4	7.8	473	4	US-09-449-249-1	Sequence 1, Appli
37	71.4	7.8	473	5	PCT-US95-10973A-25	Sequence 25, Appl
38	71.4	7.8	495	4	US-09-244-583-25	Sequence 25, Appl
39	71.4	7.8	495	4	US-09-037-983C-14	Sequence 14, Appl
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41	71.4	7.8	498	6	5194596-20	Patent No. 5194596
42	71.4	7.8	516	3	US-08-784-551C-1	Sequence 1, Appli
43	71.4	7.8	516	3	US-09-392-932-7	Sequence 7, Appli
44	71.4	7.8	516	4	US-09-574-708A-3	Sequence 3, Appli
45	71.4	7.8	516	4	US-09-037-983C-1	Sequence 1, Appli

## ALIGNMENTS

## RESULT 1

US-09-949-016-1545

; Sequence 1545, Application US/09949016

; Patent No. 6812339

; GENERAL INFORMATION:

; APPLICANT: VENTER, J. Craig et al.

; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

; TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

; FILE REFERENCE: CL001307

; CURRENT APPLICATION NUMBER: US/09/949,016

; CURRENT FILING DATE: 2000-04-14

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<a href="#">us-09-349-954a-5.rni</a>	<a href="#">Download</a>
<a href="#">us-09-349-954a-5.rnpb</a>	<a href="#">Download</a>
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<a href="#">us-09-349-954a-5.rnpn</a>	<a href="#">Download</a>
<a href="#">us-09-349-954a-7.rni</a>	<a href="#">Download</a>
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<a href="#">us-09-349-954a-9.rnpm</a>	<a href="#">Download</a>
<a href="#">us-09-349-954a-9.rnpn</a>	<a href="#">Download</a>

Comments /  
Suggestions

1	622.4	56.9	624	2	US-08-609-443B-14	Sequence 14, Appl
2	622.4	56.9	624	2	US-08-569-063C-14	Sequence 14, Appl
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5	521.6	47.7	624	2	US-08-569-063C-12	Sequence 12, Appl
6	521.6	47.7	624	4	US-08-851-896-12	Sequence 12, Appl
7	458	41.9	570	1	US-08-469-427A-10	Sequence 10, Appl
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9	458	41.9	570	2	US-08-569-063C-10	Sequence 10, Appl
10	458	41.9	570	4	US-08-851-896-10	Sequence 10, Appl
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13	359.6	32.9	565	2	US-08-569-063C-4	Sequence 4, Appli
14	359.6	32.9	565	4	US-08-851-896-4	Sequence 4, Appli
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16	332.6	30.4	591	2	US-08-609-443B-6	Sequence 6, Appli
17	332.6	30.4	591	2	US-08-569-063C-6	Sequence 6, Appli
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19	319.6	29.2	405	1	US-08-469-427A-8	Sequence 8, Appli
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26	302.8	27.7	886	4	US-08-851-896-1	Sequence 1, Appli
27	92.2	8.4	444	4	US-09-392-932-6	Sequence 6, Appli
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34	92.2	8.4	473	5	PCT-US95-10973A-25	Sequence 25, Appl
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38	92.2	8.4	516	3	US-08-784-551C-1	Sequence 1, Appli
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43	92.2	8.4	516	4	US-09-392-931-3	Sequence 3, Appli
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45	92.2	8.4	576	4	US-09-392-932-8	Sequence 8, Appli

## ALIGNMENTS

## RESULT 1

US-08-609-443B-14

; Sequence 14, Application US/08609443B

; Patent No. 5840693

; GENERAL INFORMATION:

; APPLICANT: ERIKSSON, Ulf

; APPLICANT: OLOFSSON, Birgitta

; APPLICANT: ALITALO, Kari

; APPLICANT: PAJUSOLA, Katri

; TITLE OF INVENTION: VASCULAR ENDOTHELIAL GROWTH FACTOR-B AND

; TITLE OF INVENTION: DNA CODING THEREFOR



## SCORE Search Results Details for Application 093 and Search Result us-09-349-954a-3.rnpb.

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OM nucleic - nucleic search, using sw model

Run on: March 10, 2004, 17:11:11 ; Search time 385.667 Seconds  
(without alignments)  
10374.835 Million cell updates/sec

Title: US-09-349-954A-3  
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Gapop 10.0 , Gapext 1.0

Searched: 2421054 seqs, 1828716029 residues

Total number of hits satisfying chosen parameters: 4842108

Minimum DB seq length: 0  
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Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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1	1094	100.0	1094	9	US-09-349-954A-3	Sequence 3, Appli
2	1094	100.0	1094	9	US-09-907-007-3	Sequence 3, Appli
3	1092.4	99.9	1181	14	US-10-136-819-10	Sequence 10, Appl
4	880.4	80.5	993	9	US-09-349-954A-5	Sequence 5, Appli
5	880.4	80.5	993	9	US-09-907-007-5	Sequence 5, Appli
6	876.8	80.1	1154	15	US-10-264-049-846	Sequence 846, App
7	813.4	74.4	910	9	US-09-349-954A-9	Sequence 9, Appli
8	813.4	74.4	910	9	US-09-907-007-9	Sequence 9, Appli
9	749.8	68.5	755	14	US-10-262-538-21	Sequence 21, Appl
10	749.8	68.5	755	14	US-10-007-926A-466	Sequence 466, App
11	749.8	68.5	755	15	US-10-174-128-4	Sequence 4, Appli
12	622.4	56.9	624	9	US-09-912-436-3	Sequence 3, Appli
13	622.4	56.9	624	10	US-09-961-756-14	Sequence 14, Appl
14	621.8	56.8	5695	9	US-09-912-436-10	Sequence 10, Appl
15	613.8	56.1	5695	9	US-09-912-436-9	Sequence 9, Appli
16	610.4	55.8	858	9	US-09-349-954A-7	Sequence 7, Appli
17	610.4	55.8	858	9	US-09-907-007-7	Sequence 7, Appli
18	610	55.8	663	9	US-09-244-694-19	Sequence 19, Appl
19	601	54.9	666	9	US-09-244-694-1	Sequence 1, Appli
20	595.2	54.4	1242	9	US-09-349-954A-16	Sequence 16, Appl
21	595.2	54.4	1242	9	US-09-907-007-16	Sequence 16, Appl
c 22	542.4	49.6	565	9	US-09-244-694-32	Sequence 32, Appl
23	521.6	47.7	624	10	US-09-961-756-12	Sequence 12, Appl
c 24	507.8	46.4	566	9	US-09-244-694-142	Sequence 142, App
c 25	497.2	45.4	502	9	US-09-244-694-34	Sequence 34, Appl
c 26	489.8	44.8	493	9	US-09-244-694-35	Sequence 35, Appl
c 27	467.4	42.7	469	9	US-09-244-694-39	Sequence 39, Appl
28	467.2	42.7	553	9	US-09-244-694-27	Sequence 27, Appl
c 29	461.8	42.2	490	9	US-09-244-694-37	Sequence 37, Appl
30	458	41.9	570	9	US-09-795-006A-116	Sequence 116, App
31	458	41.9	570	10	US-09-961-756-10	Sequence 10, Appl
32	456	41.7	567	9	US-09-912-436-1	Sequence 1, Appli
33	455.2	41.6	5614	9	US-09-912-436-7	Sequence 7, Appli
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35	410.4	37.5	5458	9	US-09-912-436-11	Sequence 11, Appl
36	409.8	37.5	449	10	US-09-918-995-13904	Sequence 13904, A
37	408	37.3	408	9	US-09-912-436-5	Sequence 5, Appli
38	402.4	36.8	5458	9	US-09-912-436-12	Sequence 12, Appl
39	400	36.6	445	9	US-09-244-694-22	Sequence 22, Appl
40	387.4	35.4	415	9	US-09-244-694-23	Sequence 23, Appl
41	384.8	35.2	523	9	US-09-244-694-29	Sequence 29, Appl
42	369	33.7	531	9	US-09-244-694-21	Sequence 21, Appl
43	359.6	32.9	565	10	US-09-961-756-4	Sequence 4, Appli
44	332.6	30.4	591	10	US-09-961-756-6	Sequence 6, Appli
45	323.2	29.5	374	9	US-09-244-694-33	Sequence 33, Appl

## ALIGNMENTS

RESULT 1

US-09-349-954A-3

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OM nucleic - nucleic search, using sw model

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Run on:      March 10, 2004, 17:11:11 ; Search time 109.826 Seconds
              (without alignments)
              2670.873 Million cell updates/sec
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Perfect score: 1094  
Sequence: 1 ccatgaqccctctqctccqc.....gaaggaaaaaaaaaaaaaaaaa 1094

Scoring table: IDENTITY\_NUC  
Gapop 10.0 , Gapext 1.0

Searched: 582743 seqs, 134063251 residues

Total number of hits satisfying chosen parameters: 1165486

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Post-processing: Minimum Match 0%
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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1	1094	100.0	1094	6	US-10-220-324A-1	Sequence 1, Appli
2	880.4	80.5	993	6	US-10-220-324A-3	Sequence 3, Appli
3	398.4	36.4	585	6	US-10-204-070A-3	Sequence 3, Appli
4	92.2	8.4	438	1	PCT-IL03-01085-1	Sequence 1, Appli
5	92.2	8.4	444	5	US-09-575-199C-1	Sequence 1, Appli
6	92.2	8.4	498	6	US-10-765-580-10	Sequence 10, Appl
7	92.2	8.4	541	1	PCT-US04-02188-59	Sequence 59, Appl
8	92.2	8.4	541	6	US-10-764-425-59	Sequence 59, Appl
9	92.2	8.4	576	1	PCT-US04-02974-33	Sequence 33, Appl
10	92.2	8.4	576	6	US-10-770-668-33	Sequence 33, Appl
11	92.2	8.4	1507	6	US-10-765-580-11	Sequence 11, Appl
12	92.2	8.4	4425	1	PCT-US04-05372-13	Sequence 13, Appl
13	90.6	8.3	3166	6	US-10-775-169-244	Sequence 244, App
14	79.4	7.3	438	1	PCT-IL03-01085-2	Sequence 2, Appli
15	54.6	5.0	645	6	US-10-767-701-3807	Sequence 3807, Ap
16	52.8	4.8	1347	6	US-10-767-701-11203	Sequence 11203, A
17	52	4.8	615	6	US-10-767-701-4396	Sequence 4396, Ap
18	50.2	4.6	707	6	US-10-767-701-4413	Sequence 4413, Ap
19	49.6	4.5	960	6	US-10-767-701-7244	Sequence 7244, Ap
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c 24	48.4	4.4	201	6	US-10-767-471-27003	Sequence 27003, A
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c 28	48	4.4	201	6	US-10-767-471-6787	Sequence 6787, Ap
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33	47.4	4.3	563	6	US-10-767-701-4502	Sequence 4502, Ap
c 34	47.2	4.3	1737	6	US-10-767-471-355	Sequence 355, App
c 35	47.2	4.3	11924	1	PCT-US03-31219-7	Sequence 7, Appli
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c 38	47.2	4.3	12242	6	US-10-678-816-6	Sequence 6, Appli
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44	46.4	4.2	592	6	US-10-632-150-31	Sequence 31, Appl
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## ALIGNMENTS

## RESULT 1

US-10-220-324A-1

; Sequence 1, Application US/10220324A

; GENERAL INFORMATION:

; APPLICANT: The Council of the Queensland Institute of Medical Research

; TITLE OF INVENTION: A method of treatment and prophylaxis

; FILE REFERENCE: 2386004/EJH

; CURRENT APPLICATION NUMBER: US/10/220,324A

; CURRENT FILING DATE: 2003-06-09

; PRIOR APPLICATION NUMBER: International

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11	442	44.5	591	2	US-08-569-063C-6	Sequence 6, Appli
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17	408.8	41.2	624	2	US-08-609-443B-14	Sequence 14, Appl
18	408.8	41.2	624	2	US-08-569-063C-14	Sequence 14, Appl
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20	338	34.0	624	2	US-08-609-443B-12	Sequence 12, Appl
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22	338	34.0	624	4	US-08-851-896-12	Sequence 12, Appl
23	319.6	32.2	405	1	US-08-469-427A-8	Sequence 8, Appli
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25	319.6	32.2	405	2	US-08-569-063C-8	Sequence 8, Appli
26	319.6	32.2	405	4	US-08-851-896-8	Sequence 8, Appli
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29	97.2	9.8	605	5	PCT-US95-10973A-26	Sequence 26, Appl
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33	96	9.7	576	4	US-09-392-931-5	Sequence 5, Appli
34	96	9.7	599	5	PCT-US95-10973A-87	Sequence 87, Appl
35	96	9.7	599	5	PCT-US95-10973A-89	Sequence 89, Appl
36	96	9.7	989	6	5332671-11	Patent No. 5332671
37	96	9.7	990	3	US-08-567-200A-1	Sequence 1, Appli
38	96	9.7	990	3	US-08-691-794-1	Sequence 1, Appli
39	96	9.7	990	4	US-08-882-816-1	Sequence 1, Appli
40	96	9.7	990	4	US-08-802-052B-1	Sequence 1, Appli
41	96	9.7	1299	5	PCT-US95-10973A-58	Sequence 58, Appl
42	96	9.7	1809	5	PCT-US95-10973A-79	Sequence 79, Appl
43	96	9.7	3583	4	US-09-976-594-921	Sequence 921, App
44	94.4	9.5	498	6	5194596-20	Patent No. 5194596
45	94.4	9.5	498	6	5219739-21	Patent No. 5219739

## ALIGNMENTS

## RESULT 1

US-08-469-427A-10

; Sequence 10, Application US/08469427A

; Patent No. 5607918

; GENERAL INFORMATION:

; APPLICANT: Eriksson, Ulf

; APPLICANT: Olofsson, Birgitta

; APPLICANT: Alitalo, Kari

; APPLICANT: Pajusola, Katri

; TITLE OF INVENTION: VASCULAR ENDOTHELIAL GROWTH FACTOR-B AND

; TITLE OF INVENTION: DNA CODING THEREFOR

## SCORE Search Results Details for Application 093 and Search Result us-09-349-954a-5.rnpb.

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OM nucleic - nucleic search, using sw model

Run on: March 10, 2004, 17:11:11 ; Search time 350.061 Seconds  
(without alignments)  
10374.835 Million cell updates/sec

Title: US-09-349-954A-5  
Perfect score: 993  
Sequence: 1 ccatgagccctctgctccgc.....gaaggaaaaaaaaaaaaaaaaa 993

Scoring table: IDENTITY\_NUC  
Gapop 10.0 , Gapext 1.0

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Total number of hits satisfying chosen parameters: 4842108

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Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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4	880.4	88.7	1094	9	US-09-349-954A-3	Sequence 3, Appli
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6	878.8	88.5	1181	14	US-10-136-819-10	Sequence 10, Appl
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13	565.4	56.9	567	9	US-09-912-436-1	Sequence 1, Appli
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17	536.2	54.0	755	14	US-10-262-538-21	Sequence 21, Appl
18	536.2	54.0	755	14	US-10-007-926A-466	Sequence 466, App
19	536.2	54.0	755	15	US-10-174-128-4	Sequence 4, Appli
c 20	507.8	51.1	566	9	US-09-244-694-142	Sequence 142, App
c 21	497.2	50.1	502	9	US-09-244-694-34	Sequence 34, Appl
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## ALIGNMENTS

RESULT 1

US-09-349-954A-5



# SCORE Search Results Details for Application 09349954 and Search Result us-09-349-954a-5.rnpr

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OM nucleic - nucleic search, using sw model

Run on: March 10, 2004, 17:11:11 ; Search time 3180.18 Seconds  
(without alignments)  
10986.756 Million cell updates/sec

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**SCORE Search Results Details for Application 09349954 and Search Result us-09-349-954a-5.rnpn.**

<a href="#">Score Home</a>	<a href="#">Retrieve Application</a>	<a href="#">SCORE System</a>	<a href="#">SCORE</a>	<a href="#">Comments /</a>
<a href="#">Page</a>	<a href="#">List</a>	<a href="#">Overview</a>	<a href="#">FAQ</a>	<a href="#">Suggestions</a>

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OM nucleic - nucleic search, using sw model

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Run on:      March 10, 2004, 17:11:11 ; Search time 99.6864 Seconds
              (without alignments)
              2670.873 Million cell updates/sec
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Title: US-09-349-954A-5  
Perfect score: 993  
Sequence: 1 ccatgagccctctgctccgc.....gaaggaaaaaaaaaaaaaaaaa 993

Scoring table: IDENTITY\_NUC  
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Searched: 582743 seqs, 134063251 residues

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Post-processing: Minimum Match 0%
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                  Listing first 45 summaries
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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3	511	51.5	585	6	US-10-204-070A-3	Sequence 3, Appli
4	96	9.7	498	6	US-10-765-580-10	Sequence 10, Appl
5	96	9.7	576	1	PCT-US04-02974-33	Sequence 33, Appl
6	96	9.7	576	6	US-10-770-668-33	Sequence 33, Appl
7	96	9.7	1507	6	US-10-765-580-11	Sequence 11, Appl
8	96	9.7	4425	1	PCT-US04-05372-13	Sequence 13, Appl
9	94.4	9.5	3166	6	US-10-775-169-244	Sequence 244, App
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14	79.4	8.0	438	1	PCT-IL03-01085-2	Sequence 2, Appli
15	45.8	4.6	707	6	US-10-767-701-4413	Sequence 4413, Ap
c 16	42.6	4.3	1220	6	US-10-767-701-13830	Sequence 13830, A
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21	41.8	4.2	1721	6	US-10-775-920-88	Sequence 88, Appl
22	41.8	4.2	2238	6	US-10-775-920-87	Sequence 87, Appl
23	41	4.1	453	6	US-10-767-701-26919	Sequence 26919, A
c 24	41	4.1	1424	6	US-10-778-859-75	Sequence 75, Appl
25	41	4.1	1428	6	US-10-778-859-20	Sequence 20, Appl
26	41	4.1	1799	6	US-10-778-859-19	Sequence 19, Appl
27	41	4.1	1804	6	US-10-775-920-84	Sequence 84, Appl
28	41	4.1	8181	6	US-10-778-859-18	Sequence 18, Appl
c 29	40.6	4.1	1355	6	US-10-045-674A-593	Sequence 593, App
c 30	40.4	4.1	636	6	US-10-767-701-24914	Sequence 24914, A
c 31	39.8	4.0	918	6	US-10-767-701-15319	Sequence 15319, A
c 32	39.8	4.0	965	6	US-10-767-701-15468	Sequence 15468, A
33	39.4	4.0	1347	6	US-10-767-701-11203	Sequence 11203, A
c 34	39.2	3.9	1588	6	US-10-767-701-14522	Sequence 14522, A
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c 37	38.4	3.9	2269	5	US-09-860-298B-1	Sequence 1, Appli
c 38	38.4	3.9	4739	6	US-10-770-726-2	Sequence 2, Appli
39	38.2	3.8	78568	6	US-10-767-471-10897	Sequence 10897, A
40	38	3.8	604	6	US-10-767-701-7783	Sequence 7783, Ap
41	38	3.8	659	6	US-10-767-701-4135	Sequence 4135, Ap
42	38	3.8	3917	6	US-10-775-920-127	Sequence 127, App
c 43	37.8	3.8	965	1	PCT-US04-05654-2095	Sequence 2095, Ap
44	37.8	3.8	2571	5	US-09-999-183A-4	Sequence 4, Appli
45	37.8	3.8	3286	6	US-10-767-471-162	Sequence 162, App

## ALIGNMENTS

## RESULT 1

US-10-220-324A-3

; Sequence 3, Application US/10220324A

; GENERAL INFORMATION:

; APPLICANT: The Council of the Queensland Institute of Medical Research

; TITLE OF INVENTION: A method of treatment and prophylaxis

; FILE REFERENCE: 2386004/EJH

; CURRENT APPLICATION NUMBER: US/10/220,324A

; CURRENT FILING DATE: 2003-06-09

; PRIOR APPLICATION NUMBER: International



1	411.2	47.9	570	1	US-08-469-427A-10	Sequence 10, Appl
2	411.2	47.9	570	2	US-08-609-443B-10	Sequence 10, Appl
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4	411.2	47.9	570	4	US-08-851-896-10	Sequence 10, Appl
5	409.4	47.7	624	2	US-08-609-443B-14	Sequence 14, Appl
6	409.4	47.7	624	2	US-08-569-063C-14	Sequence 14, Appl
7	409.4	47.7	624	4	US-08-851-896-14	Sequence 14, Appl
8	339	39.5	624	2	US-08-609-443B-12	Sequence 12, Appl
9	339	39.5	624	2	US-08-569-063C-12	Sequence 12, Appl
10	339	39.5	624	4	US-08-851-896-12	Sequence 12, Appl
11	338.4	39.4	565	1	US-08-469-427A-4	Sequence 4, Appli
12	338.4	39.4	565	2	US-08-609-443B-4	Sequence 4, Appli
13	338.4	39.4	565	2	US-08-569-063C-4	Sequence 4, Appli
14	338.4	39.4	565	4	US-08-851-896-4	Sequence 4, Appli
15	319.6	37.2	405	1	US-08-469-427A-8	Sequence 8, Appli
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18	319.6	37.2	405	4	US-08-851-896-8	Sequence 8, Appli
19	311.4	36.3	591	1	US-08-469-427A-6	Sequence 6, Appli
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22	311.4	36.3	591	4	US-08-851-896-6	Sequence 6, Appli
23	253	29.5	886	1	US-08-469-427A-1	Sequence 1, Appli
24	253	29.5	886	2	US-08-609-443B-1	Sequence 1, Appli
25	253	29.5	886	2	US-08-569-063C-1	Sequence 1, Appli
26	253	29.5	886	4	US-08-851-896-1	Sequence 1, Appli
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36	92.2	10.7	495	4	US-09-037-983C-14	Sequence 14, Appl
37	92.2	10.7	498	6	5194596-20	Patent No. 5194596
38	92.2	10.7	516	3	US-08-784-551C-1	Sequence 1, Appli
39	92.2	10.7	516	4	US-09-392-932-7	Sequence 7, Appli
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42	92.2	10.7	516	4	US-09-428-909A-1	Sequence 1, Appli
43	92.2	10.7	516	4	US-09-392-931-3	Sequence 3, Appli
44	92.2	10.7	545	4	US-09-244-583-1	Sequence 1, Appli
45	92.2	10.7	576	4	US-09-392-932-8	Sequence 8, Appli

## ALIGNMENTS

## RESULT 1

US-08-469-427A-10

; Sequence 10, Application US/08469427A

; Patent No. 5607918

; GENERAL INFORMATION:

; APPLICANT: Eriksson, Ulf

; APPLICANT: Olofsson, Birgitta

; APPLICANT: Alitalo, Kari

; APPLICANT: Pajusola, Katri

; TITLE OF INVENTION: VASCULAR ENDOTHELIAL GROWTH FACTOR-B AND

; TITLE OF INVENTION: DNA CODING THEREFOR

## SCORE Search Results Details for Application 09349954 and Search Result us-09-349-954a-7.rnpb.

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This page gives you Search Results detail for the Application 09349954 and Search Result us-09-349-954a-7.rnpb.

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OM nucleic - nucleic search, using sw model

Run on: March 10, 2004, 17:11:11 ; Search time 302.47 Seconds  
(without alignments)  
10374.835 Million cell updates/sec

Title: US-09-349-954A-7  
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Gapop 10.0 , Gapext 1.0

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Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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2	858	100.0	858	9	US-09-907-007-7	Sequence 7, Appli
3	713	83.1	993	9	US-09-349-954A-5	Sequence 5, Appli
4	713	83.1	993	9	US-09-907-007-5	Sequence 5, Appli
5	707.8	82.5	1154	15	US-10-264-049-846	Sequence 846, App
6	610.4	71.1	1094	9	US-09-349-954A-3	Sequence 3, Appli
7	610.4	71.1	1094	9	US-09-907-007-3	Sequence 3, Appli
8	608.8	71.0	1181	14	US-10-136-819-10	Sequence 10, Appl
9	447.4	52.1	910	9	US-09-349-954A-9	Sequence 9, Appli
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c 11	440.4	51.3	565	9	US-09-244-694-32	Sequence 32, Appl
c 12	438.4	51.1	469	9	US-09-244-694-39	Sequence 39, Appl
c 13	435.8	50.8	493	9	US-09-244-694-35	Sequence 35, Appl
c 14	435.8	50.8	502	9	US-09-244-694-34	Sequence 34, Appl
c 15	421.6	49.1	566	9	US-09-244-694-142	Sequence 142, App
c 16	415.6	48.4	490	9	US-09-244-694-37	Sequence 37, Appl
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22	411.2	47.9	570	10	US-09-961-756-10	Sequence 10, Appl
23	411.2	47.9	5614	9	US-09-912-436-7	Sequence 7, Appli
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## ALIGNMENTS

RESULT 1

US-09-349-954A-7



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# SCORE Search Results Details for Application 09349954 and Search Result us-09-349-954a-7.rnnpn.

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OM nucleic - nucleic search, using sw model

Run on: March 10, 2004, 17:11:11 ; Search time 86.1339 Seconds  
(without alignments)  
2670.873 Million cell updates/sec

Title: US-09-349-954A-7  
Perfect score: 858  
Sequence: 1 ccatgagccctctgctcgc.....gaaggaaaaaaaaaaaaaaaaa 858

Scoring table: IDENTITY\_NUC  
Gapop 10.0 , Gapext 1.0

Searched: 582743 seqs, 134063251 residues

Total number of hits satisfying chosen parameters: 1165486

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Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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4	92.2	10.7	438	1	PCT-IL03-01085-1	Sequence 1, Appli
5	92.2	10.7	444	5	US-09-575-199C-1	Sequence 1, Appli
6	92.2	10.7	498	6	US-10-765-580-10	Sequence 10, Appl
7	92.2	10.7	541	1	PCT-US04-02188-59	Sequence 59, Appl
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13	90.6	10.6	3166	6	US-10-775-169-244	Sequence 244, App
14	79.4	9.3	438	1	PCT-IL03-01085-2	Sequence 2, Appli
15	41	4.8	453	6	US-10-767-701-26919	Sequence 26919, A
16	38.2	4.5	78568	6	US-10-767-471-10897	Sequence 10897, A
17	37.8	4.4	2571	5	US-09-999-183A-4	Sequence 4, Appli
18	36.6	4.3	1458	6	US-10-767-701-10527	Sequence 10527, A
19	36	4.2	2488	1	PCT-US04-06020-31	Sequence 31, Appl
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## ALIGNMENTS

## RESULT 1

US-10-220-324A-3

; Sequence 3, Application US/10220324A

; GENERAL INFORMATION:

; APPLICANT: The Council of the Queensland Institute of Medical Research

; TITLE OF INVENTION: A method of treatment and prophylaxis

; FILE REFERENCE: 2386004/EJH

; CURRENT APPLICATION NUMBER: US/10/220,324A

; CURRENT FILING DATE: 2003-06-09

; PRIOR APPLICATION NUMBER: International

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## ALIGNMENTS

## RESULT 1

US-08-609-443B-14

; Sequence 14, Application US/08609443B

; Patent No. 5840693

; GENERAL INFORMATION:

; APPLICANT: ERIKSSON, Ulf

; APPLICANT: OLOFSSON, Birgitta

; APPLICANT: ALITALO, Kari

; APPLICANT: PAJUSOLA, Katri

; TITLE OF INVENTION: VASCULAR ENDOTHELIAL GROWTH FACTOR-B AND

; TITLE OF INVENTION: DNA CODING THEREFOR

## SCORE Search Results Details for Application 093 and Search Result us-09-349-954a-9.rnpb.

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OM nucleic - nucleic search, using sw model

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## ALIGNMENTS

RESULT 1

US-09-349-954A-9



**SCORE Search Results Details for Application 09349954 and Search Result us-09-349-954a-9.rnpm.**

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OM nucleic - nucleic search, using sw model

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4: /cgn2\_6/ptodata/2/pna/US08\_NEW\_COMB.seq:\*  
5: /cgn2\_6/ptodata/2/pna/US09\_NEW\_COMB.seq:\*  
6: /cgn2\_6/ptodata/2/pna/US10\_NEW\_COMB.seq:\*  
7: /cgn2\_6/ptodata/2/pna/US60\_NEW\_COMB.seq:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.	Score	Match	Length	DB ID	Description
		%			
		Query			

1	813.4	89.4	1094	6	US-10-220-324A-1	Sequence 1, Appli
2	706.6	77.6	993	6	US-10-220-324A-3	Sequence 3, Appli
3	347.2	38.2	585	6	US-10-204-070A-3	Sequence 3, Appli
4	71.4	7.8	438	1	PCT-IL03-01085-1	Sequence 1, Appli
5	71.4	7.8	444	5	US-09-575-199C-1	Sequence 1, Appli
6	71.4	7.8	498	6	US-10-765-580-10	Sequence 10, Appl
7	71.4	7.8	541	1	PCT-US04-02188-59	Sequence 59, Appl
8	71.4	7.8	541	6	US-10-764-425-59	Sequence 59, Appl
9	71.4	7.8	576	1	PCT-US04-02974-33	Sequence 33, Appl
10	71.4	7.8	576	6	US-10-770-668-33	Sequence 33, Appl
11	71.4	7.8	1507	6	US-10-765-580-11	Sequence 11, Appl
12	71.4	7.8	4425	1	PCT-US04-05372-13	Sequence 13, Appl
13	69.8	7.7	3166	6	US-10-775-169-244	Sequence 244, App
14	58.6	6.4	438	1	PCT-IL03-01085-2	Sequence 2, Appli
15	54.6	6.0	645	6	US-10-767-701-3807	Sequence 3807, Ap
16	52.8	5.8	1347	6	US-10-767-701-11203	Sequence 11203, A
17	52	5.7	615	6	US-10-767-701-4396	Sequence 4396, Ap
18	50.2	5.5	707	6	US-10-767-701-4413	Sequence 4413, Ap
19	49.6	5.5	960	6	US-10-767-701-7244	Sequence 7244, Ap
c 20	49	5.4	1000	6	US-10-779-543-8571	Sequence 8571, Ap
c 21	49	5.4	3195	1	PCT-US03-35026-54	Sequence 54, Appl
22	48.8	5.4	584	6	US-10-767-701-4186	Sequence 4186, Ap
23	48.6	5.3	1092	6	US-10-767-701-9739	Sequence 9739, Ap
c 24	48.4	5.3	201	6	US-10-767-471-6782	Sequence 6782, Ap
c 25	48.4	5.3	201	6	US-10-767-471-27003	Sequence 27003, A
26	48.4	5.3	667	6	US-10-767-701-8805	Sequence 8805, Ap
27	48.4	5.3	1337	6	US-10-779-543-5579	Sequence 5579, Ap
c 28	48	5.3	201	6	US-10-767-471-6785	Sequence 6785, Ap
c 29	48	5.3	201	6	US-10-767-471-6787	Sequence 6787, Ap
c 30	48	5.3	201	6	US-10-767-471-27022	Sequence 27022, A
c 31	48	5.3	201	6	US-10-767-471-27024	Sequence 27024, A
32	47.8	5.3	1388	6	US-10-767-701-14298	Sequence 14298, A
33	47.6	5.2	1215	6	US-10-767-701-8767	Sequence 8767, Ap
34	47.4	5.2	563	6	US-10-767-701-4502	Sequence 4502, Ap
c 35	47.2	5.2	1737	6	US-10-767-471-355	Sequence 355, App
c 36	47.2	5.2	11924	1	PCT-US03-31219-7	Sequence 7, Appli
c 37	47.2	5.2	11924	6	US-10-678-816-7	Sequence 7, Appli
c 38	47.2	5.2	12242	1	PCT-US03-31219-6	Sequence 6, Appli
c 39	47.2	5.2	12242	6	US-10-678-816-6	Sequence 6, Appli
c 40	47.2	5.2	24260	6	US-10-767-471-10667	Sequence 10667, A
41	47	5.2	1021	6	US-10-767-701-11876	Sequence 11876, A
42	46.8	5.1	573	6	US-10-767-701-4301	Sequence 4301, Ap
43	46.8	5.1	797	6	US-10-767-701-4259	Sequence 4259, Ap
44	46.8	5.1	862	6	US-10-767-701-10425	Sequence 10425, A
45	46.4	5.1	592	6	US-10-632-150-31	Sequence 31, Appl

## ALIGNMENTS

## RESULT 1

US-10-220-324A-1

; Sequence 1, Application US/10220324A

; GENERAL INFORMATION:

; APPLICANT: The Council of the Queensland Institute of Medical Research

; TITLE OF INVENTION: A method of treatment and prophylaxis

; FILE REFERENCE: 2386004/EJH

; CURRENT APPLICATION NUMBER: US/10/220,324A

; CURRENT FILING DATE: 2003-06-09

; PRIOR APPLICATION NUMBER: International